

APPROVED BY DRAFTSMAN	Q.G. FIG.	
	CLASS	SUBCLASS

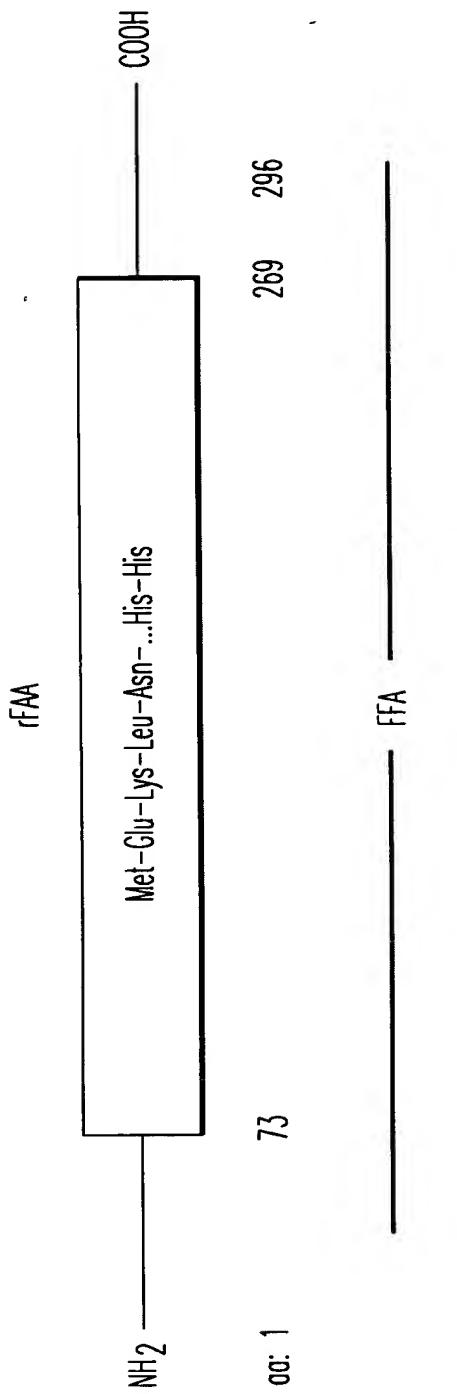


FIG. 1

APPROVED BY DRAFTSMAN	O.G. FIG.
	CLASS SUBCLASS

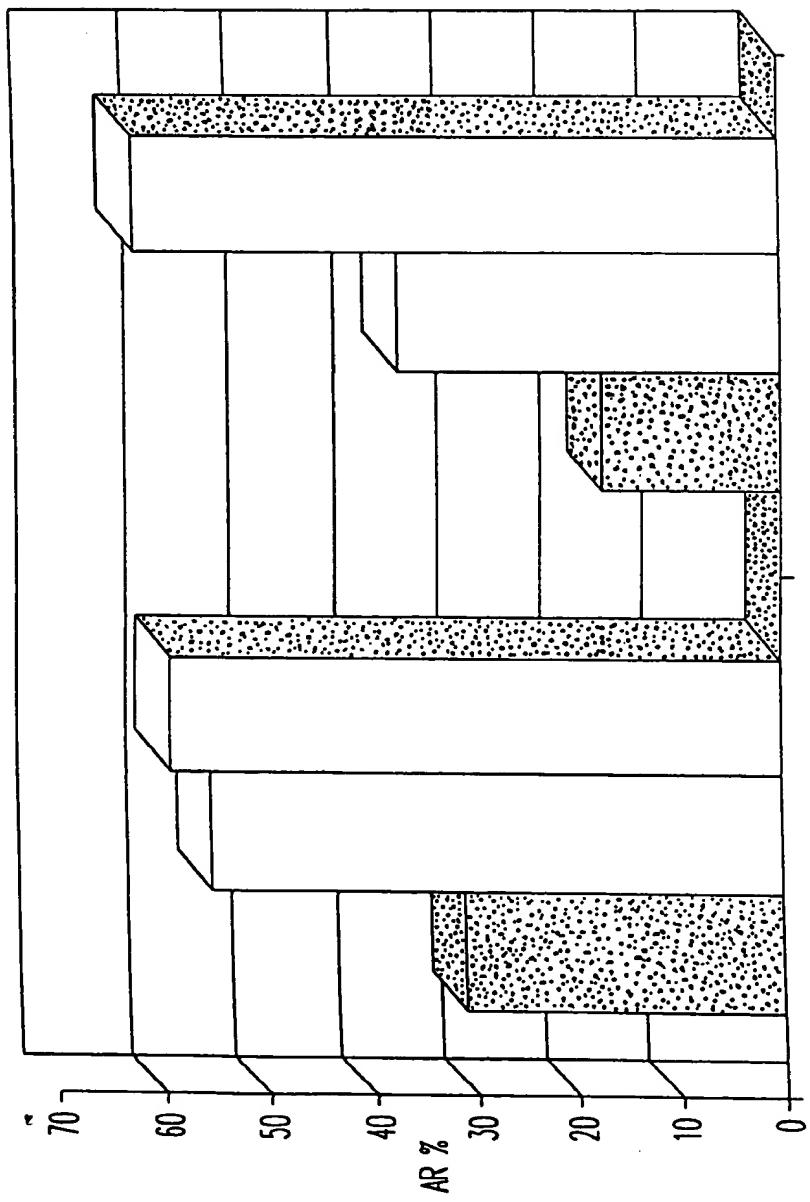
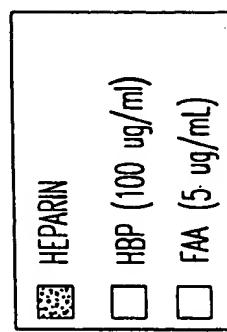


FIG. 2

FAA NEGATIVE
FAA PRESENT

APPROVED BY DRAFTSMAN	O.G. FIG.	
	CLASS	SUBCLASS

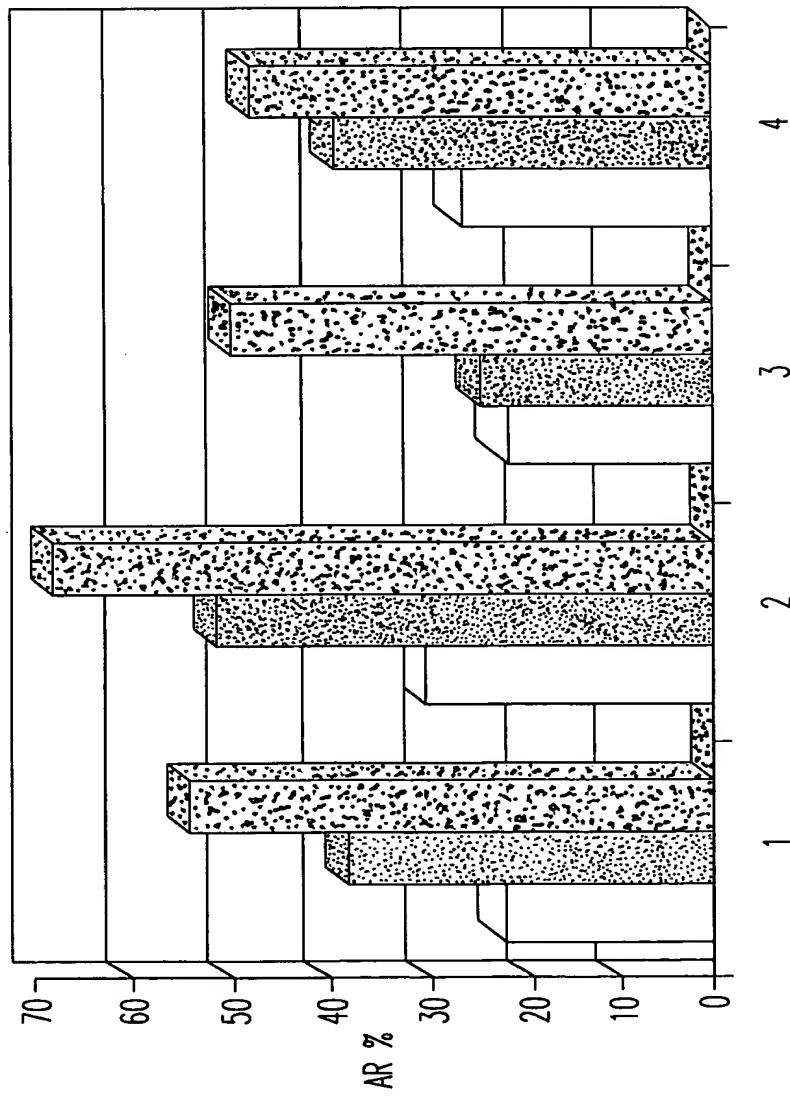
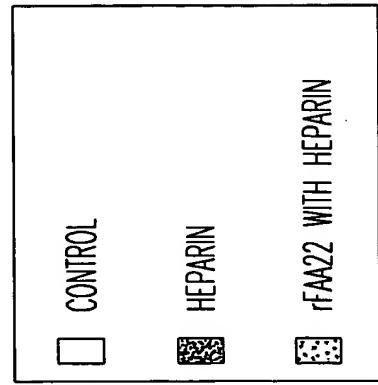


FIG. 3

BULL

4

3

2

1

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O.G. FIG.

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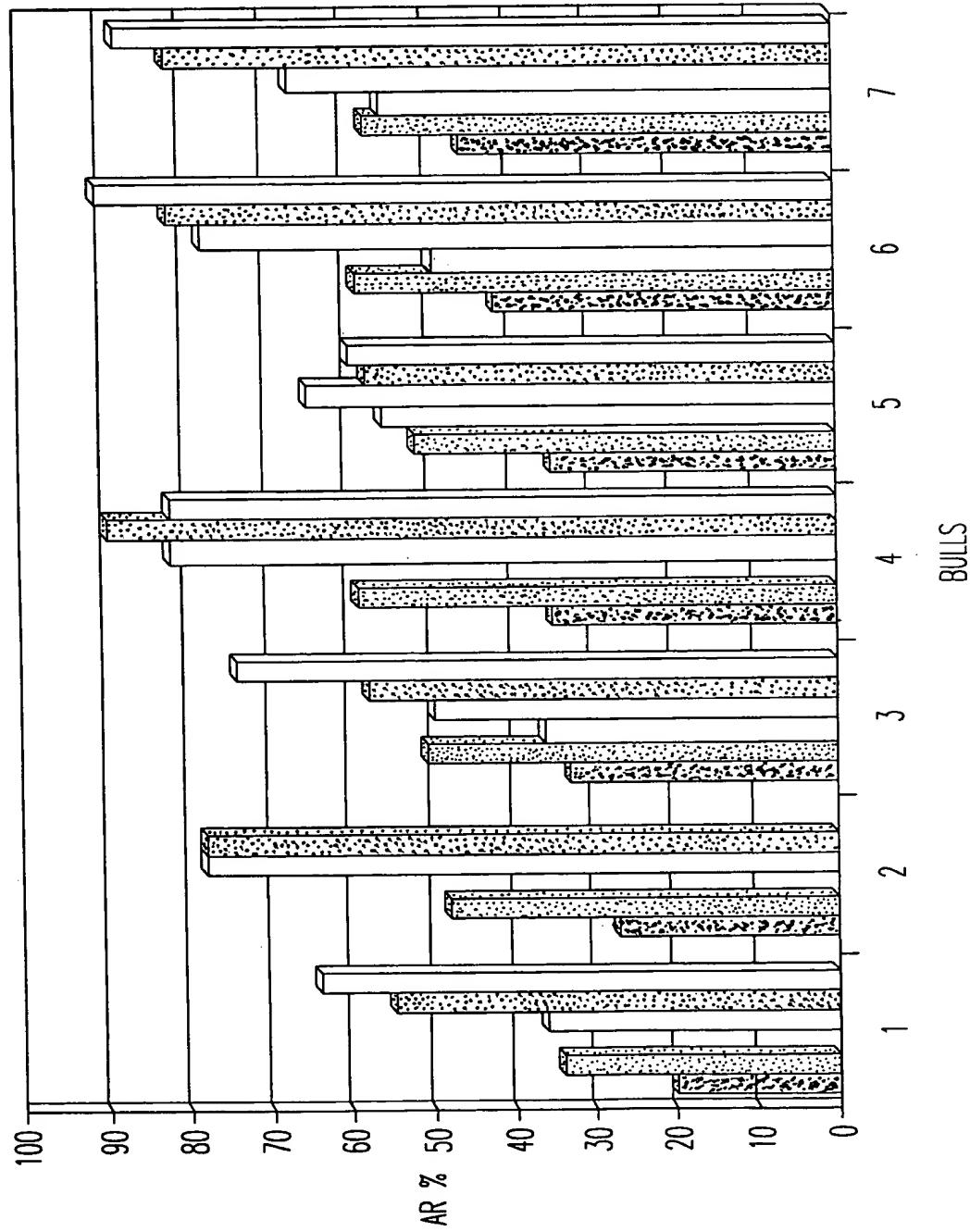
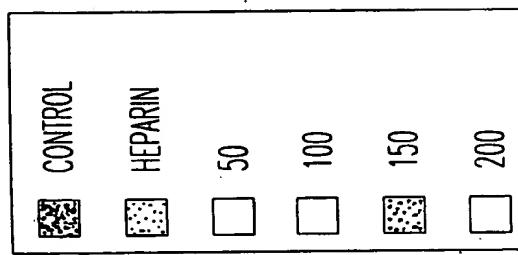


FIG. 4

APPROVED BY DRAFTSMAN	O.G. FIG.	
	CLASS	SUBCLASS



1 ACAACAGGAT CTGCCCCATA CTGATGGAGA AGCTAACCGG AAATTCAAGA
 51 AAAGGCATAA CATAACAATA TGTGATTAGC TCTCGCCTTG GAAGAAACAC
 101 ATATAAAGAA CAGTATGCCT TTCTCTATAA AGAAAAGCTA GTGTCTGTA
 151 AACAAAGCTA CCTCTACAC GACTATCAGG CTGGAGACGC AGATGTGTTT
 201 TCCAGGGAAC CCTTTGTGGT CTGGTTCCAG TCACCCCTACA CCGCTGTCAA
 251 GGACTTCGTG ATTGTCCCCC TGACACACCAC CCCTGAGACA TCCGTTAGAG
 301 AGATTGATGA GCTGGCTGAT GTCTACACAG ATGTGAAACG TCGCTGGAAT
 351 GCAGAGAATT TCATTTCAT GGGTGACTTC AATGCTGGCT GCAGCTACGT
 401 CCCCAAGAAG GCCTGGAAGG ACATCCGCCT GAGGACGGAC CCCAAGTTG
 451 TTTGGCTGAT CGGGGACCAA GAGGACACCA CGGTCAAGAA GAGCACAAAC
 501 TGCGCCTATG ACAGGATCGT GCTTAGAGGA CAAAATATTG TCAACTCTGG
 551 TGGTCCTCAA TCAAACCTCG TCTTGATTT CCAGAAAGCT TACAGGTTGT
 601 CTGAATCGAA GGCCCTGGAT GTCAGCGACC ACTTTCCAGT TCATCATCAT
 651 CATCATCATG AAGAACCATG A

FIG. 5

TOP SECRET - INTELLIGENCE

5' GAGAAGCTAACCGGAAATTCAAGAAAAGGCATAACATACAACATGTGATTAGCTCTGGC
 1 E K L N G N S R K G I T Y N Y V I S S R
 a CTTGGAAGAAACACATATAAAGAACAGTATGCCTTCTCTATAAAGAAAAGCTAGTGTCT
 61 L G R N T Y K E Q Y A F L Y K E K L V S -
 a GTAAAACAAAGCTACCTCTACCACGACTATCAGGCTGGAGACGCAGATGTGTTTCCAGG
 121 V K Q S Y L Y H D Y Q A G D A D V F S R -
 a GAACCTTTGTGGCTGGTCCAGTCACCCCTACACCGCTGTCAAGGACTTCGTGATTGTC
 181 E P F V V W F Q S P Y T A V K D F V I V -
 a CCCCTGCACACCACCCCTGAGACATCCGTTAGAGAGATTGATGAGCTGGCTGATGTCTAC
 241 P L H T T P E T S V R E I D E L A D V Y -
 a ACAGATGTGAAACGTCGCTGGAATGCAGAGAATTTCATTTCATGGGTACTTCATGCT
 301 T D V K R R W N A E N F I F M G D F N A -
 a GGCTGCAGCTACGTCCCCAAGAAGGCCTGGAGGACATCCGCTGAGGACGGACCCCAAG
 361 G C S Y V P K K A W K D I R L R T D P K -
 a TTCTGTTGGCTGATGGGGACCAAGAGGACACCACGGTCAAGAAGAGGACACAAACTGCGCC
 421 F V W L I G D Q E D T T V K K S T N C A -
 a TATGACAGGATCGTGCTTAGAGGACAAAATATTGTCAACTCTGGTGGTCCTCAATCAAAC
 481 Y D R I V L R G Q N I V N S G G P Q S N -
 a CTCGTCTTGTATTCCAGAAAGCTTACAGGTTGTCTGAATCGAAGGCCCTGG
 541 L V F D F Q K A Y R L S E S K A L - 592 3'
 a